

1636 1600

CRF Errors Corrected by the STIC Syst

Branch

Serial Number:

09/621,781

CRF Processing Date:

11/9/01

Edited by:

DC

Verified by:

(STIC sta

☐

Changed a file from non-ASCII to ASCII

☐

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐

Edited a formal error in the Current Application Data section, specifically:

ENTERED

#7

☐

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____

☐

Added the mandatory heading and subheadings for "Current Application Data".

☐

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐

Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐

Inserted colons after headings/subheadings. Headings edited included: _____

☐

Deleted extra, invalid, headings used by an applicant, specifically: _____

RECEIVED

NOV 29 2001

TECH CENTER 1600/2900

☒

Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____

☐

Inserted mandatory headings, specifically: _____

☐

Corrected an obvious error in the response, specifically: _____

☐

Edited identifiers where upper case is used but lower case is required, or vice versa.

☐

Corrected an error in the Number of Sequences field, specifically: _____

☐

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐

Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐

Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/621,781

DATE: 11/09/2001

TIME: 15:58:35

Input Set : A:\#351657 v1 - A34690 - Sequence Listing.txt

Output Set: N:\CRF3\11092001\I621781.raw

**Does Not Comply
Corrected Diskette Needed**

4 <110> APPLICANT: Fisher, Paul
5 Su, Zao-Zhong
7 <120> TITLE OF INVENTION: Nucleic Acids Comprising Regions of the
8 Rat PEG-3 Promoter that Display Elevated Expression in Human
9 Cancer Cells and Uses Thereof
12 <130> FILE REFERENCE: A34690
14 <140> CURRENT APPLICATION NUMBER: 09/621,781
15 <141> CURRENT FILING DATE: 2000-07-21
17 <160> NUMBER OF SEQ ID NOS: 12
19 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

171 <210> SEQ ID NO: 12
172 <211> LENGTH: 20
173 <212> TYPE: DNA
174 <213> ORGANISM: Artificial Sequence
176 <220> FEATURE:
177 <223> OTHER INFORMATION: synthetic oligonucleotide
179 <400> SEQUENCE: 12
180 cacaacaagg gtagagaggt

20

E--> 181 ny02:351657.1

delete

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/621,781

DATE: 11/09/2001

TIME: 15:58:36

Input Set : A:\#351657 v1 - A34690 - Sequence Listing.txt

Output Set: N:\CRF3\11092001\I621781.raw

L:181 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:12
L:181 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:12
L:181 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:181 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:24 SEQ:12
L:181 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
L:181 M:112 C: (48) String data converted to lower case,
L:181 M:252 E: No. of Seq. differs, <211>LENGTH:Input:20 Found:24 SEQ:12